

ID HPCCPA standard; genomic RNA; VRL; 411 BP.
 XX
 AC D30046;
 XX
 SV D30046.1
 XX
 DT 13-MAY-1994 (Rel. 39, Created)
 DT 07-JUL-2003 (Rel. 76, Last updated, Version 10)
 XX
 DE Hepatitis C virus (individual_isolate Td-3/93) gene for polyprotein
 DE precursor, partial cds (core protein (carboxy terminus) and E1 envelope
 DE protein (amino terminus half)).
 XX
 KW core protein; E1 envelope protein.
 XX
 OS Hepatitis C virus
 OC Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
 OC Hepacivirus.
 XX
 RN [1]
 RP 1-411
 RA Hotta H.;
 RT ;
 RL Submitted (28-APR-1994) to the EMBL/GenBank/DDBJ databases.
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 RL 7-5-1 Kusunoki-cho, Chuo-ku, Kobe, Hyogo 650, Japan
 RL (Tel:078-341-7451(ex.3301), Fax:078-351-6347)
 XX
 RN [2]
 RP 1-411
 RA Hotta H.;
 RT ;
 RL Unpublished.
 XX
 RN [3]
 RX MEDLINE; 95189942.
 RX PUBMED; 7883898.
 RA Hotta H., Handajani R., Ingelusida M., Soemarto W., Doi H., Miyajima H.,
 RA Homma M.;
 RT "Subtype analysis of hepatitis C virus in Indonesia on the basis of NS5b
 RT region sequences";
 RL J. Clin. Microbiol. 32(12):3049-3051(1994).
 XX
 DR GOA; Q81293.
 DR SPTREMBL; Q81293; Q81293.
 XX
 CC Submitted (28-Apr-1994) to DDBJ by:
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 FH Key Location/Qualifiers
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 FT /organism="Hepatitis C virus"
 FT /isolate="Td-3/93"
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FT /db_xref="SPTREMBL:Q81293"
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FT amino terminal half of the E1 envelope protein of hepatitis
FT C virus"
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FT /translation="LSCLLTPTAGLEYRNASGLYIVTNDCSNSSTIVYEAQDIILHMPGC
FT DPCVRSNGNTRCWTVPVSSTVAVGRAGAATASLRTHVDMMVGAATLCSALYVGDLCGALF
FT LVGQGFSWRHRQHWTVQDCNCSIYPGHLTGHRM"
XX

SQ Sequence 411 BP; 79 A; 120 C; 108 G; 104 T; 0 other;
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cacatgcctg gctgtgaccc ctgcgtacgc tctggcaaca catcaagggtg ctggacocct 180
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ggagcgcttt ttcttggttg gcaggggttc tcatggagac atcgccagca ttggactgtc 360
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